

GenCore version 5.1.4\_p5\_4578  
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score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Pred. No. is the number of results predicted by chance to have a

66	24.8	0.5	43	€	AX395321	Sequence	AX441075	Sequence	
67	24.8	0.5	43	€	AX443022	Sequence	AX187454	Sequence	
c	6.9	24.8	0.5	43	€	AY459616	Sequence	AP082115	Sequence
c	7.0	24.8	0.5	46	€	AX287578	Sequence	AX164808	Sequence
c	71	24.8	0.5	46	€	AX287582	Sequence	AR174581	Sequence
c	72	24.6	0.5	44	€	AP039858	Sequence	AX106717	Sequence
c	73	24.6	0.5	44	€	ARN1540	Sequence	179494	Sequence 1
c	74	24.6	0.5	48	€	AP021989	Sequence	A08914	Sequence
c	75	24.6	0.5	48	€	AP043404	Sequence	A08914	Sequence
c	76	24.6	0.5	48	€	AP062319	Sequence	AX430213	Sequence
c	77	24.4	0.5	30	€	AP181778	Sequence	AF029415	Sequence
c	78	24.4	0.5	35	€	AY079109	Sequence	AP1010128	Sequence
c	79	24.4	0.5	35	€	AP029410	Sequence	TG29924	Sequence 37
c	80	24.4	0.5	43	€	129930	Sequence	AX315321	Sequence
c	81	24.4	0.5	43	€	178446	Sequence	AX443C22	Sequence
c	82	24.4	0.5	45	€	AX287571	Sequence	AX459616	Sequence
c	83	24.4	0.5	45	€	AX287575	Sequence	178847	Sequence 2
c	84	24.4	0.5	46	€	AX287578	Sequence	178649	Sequence 4
c	85	24.4	0.5	46	€	AX287582	Sequence	178851	Sequence 6
c	86	24.4	0.5	50	€	AX157157	Sequence	178856	Sequence 11
c	87	24.2	0.4	32	€	AP092289	Sequence	AX206881	Sequence
c	88	24.2	0.4	32	€	AP05340	Sequence	AR003731	Sequence
c	89	24.2	0.4	32	€	AY099615	Sequence	AP01067	Sequence
c	90	24.2	0.4	32	€	AP120128	Sequence	AP05273	Sequence
c	91	24.2	0.4	37	€	129931	Sequence	AR141202	Sequence
c	92	24.2	0.4	37	€	E50166	Vector expr	AR141439	Sequence
c	93	24.2	0.4	45	€	AP021973	Sequence	11911	Sequence 21
c	94	24.2	0.4	45	€	AX172448	Sequence	140384	Sequence 21
c	95	24.2	0.4	45	€	AY287569	Sequence	AX43967	Sequence
c	96	24.2	0.4	45	€	AX287573	Sequence	AX194782	Sequence
c	97	24.2	0.4	47	€	AP038056	Sequence	AX13326	Sequence
c	98	24.2	0.4	24	€	AP09847	Sequence	AR082114	Sequence
c	99	24.2	0.4	31	€	BD011883	Detection	AY082994	Sequence
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101	24.4	0.4	31	€	AY099603	Sequence	AX052919	Sequence	
102	24.4	0.4	38	€	AR039603	Sequence	AX333685	Sequence	
103	24.4	0.4	42	€	AP039671	Sequence	AR157852	Sequence	
104	24.4	0.4	42	€	AP043386	Sequence	AF162080	Sequence	
105	24.4	0.4	42	€	AP204721	Sequence	AR1626605	Sequence	
106	24.4	0.4	42	€	AP021301	Sequence	AX04808	Sequence	
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c	112	24.4	0.4	46	€	AX287579	Sequence	BD2995	Sequence 3
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c	119	23.8	0.4	50	€	AX158156	Sequence	AX104903	Sequence
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C 218	23	0.4	6	AR202472	Sequence	AF178102	Sequence
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C	360	22	6	AX418163	Sequence AX418163
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379	22	6	I72101	Sequence I72101	
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381	22	6	AX297738	Sequence AX297738	
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385	22	6	G73658	Sequence G73658	
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389	21	6	AP09847	Sequence AP09847	
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391	21	6	AP094848	Sequence AP094848	
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C	402	21	6	AX480612	Sequence AX480612
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C	404	21	6	AX30501	Sequence AX30501
405	21	6	I32114	Sequence I32114	
C	406	21	6	A65286	Sequence A65286
407	21	6	AR116106	Sequence AR116106	
C	408	21	6	AR150434	Sequence AR150434
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411	21	6	A65285	Sequence A65285	
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413	21	6	AR172429	Sequence AR172429	
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C	416	21	6	AP09768	Sequence AP09768
417	21	6	T89907	Sequence T89907	
C	418	21	6	AB4756	Sequence AB4756
419	21	6	AP053158	Sequence AP053158	
C	420	21	6	AR071990	Sequence AR071990
421	21	6	AP093188	Sequence AP093188	
C	422	21	6	AR10948	Sequence AR10948
423	21	6	AP123332	Sequence AP123332	
C	424	21	6	AX157184	Sequence AX157184
425	21	6	AX163980	Sequence AX163980	
C	426	21	6	AX163446	Sequence AX163446
427	21	6	E115733	Sequence E115733	
C	428	21	6	E15768	Sequence E15768
429	21	6	E30813	Novel prpte	
C	430	21	6	E54866	Process for

5	0	6	21.8	C-4	331	I47698	Sequence 11
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5	0	6	21.6	C-35	AR08537	AP084537	Sequence
5	0	6	21.6	C-36	AR064078	AP064078	Sequence
5	0	6	21.6	C-37	BD008477	B0D08477	Targeting
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5	0	6	21.6	C-39	AP064077	AP064077	Sequence
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VERSION					synthetic construct.		
KEYWORDS					artificial sequences.		
SOURCE	Unknown				PREFERENCE		
ORGANISM	Unclassified				1 (bases 1 to 37)		
REFERENCE	1 (bases 1 to 30)				Authors		
AUTHORS	Dawkins R Letts and Abraham J. Joseph				Plano, L., Bernad Miana, A., Dominguez Lopez, G. and Garcia Piaz, M.		
TITLE	Method for determining ancestral haplotypes using haplospecific geometric elements within the major histocompatibility complex multigene cluster				Title		
JOURNAL	Patent. US 6383747-A 4 JUN 2002,				Journal		
FEATURES	/organism="unknown"				Patent	WO 012542 A 25-12-APP 2001;	
BASE COUNT	0 a 15 c 0 g				CONFIRM.	SUPERIOR INVESTIGATIONS SCIENTIFICAS (ES)	
ORIGIN	1..30				Location/Qualifiers		
FEATURES	/organism="unknown"				source	1..37	
BASE COUNT	0 a 15 c 0 g				/organism="synthetic construct"		
ORIGIN	1..30				/db_xref="taxon:32630"		
Query Match					/note="poly dT"		
Best Local Similarity	83.3%						
Matches	30;						
DEFINITION	Conservative						
ACCESION	129931						
VERSION	GI:1820722						
KEYWORDS							
SOURCE	Unknown						
ORGANISM	Unclassified						
REFERENCE	1 (bases 1 to 37)						
AUTHORS	Pickup, D.J., Patel, D. and Antczak, J.R.						
TITLE	Site-specific RNA cleavage						
JOURNAL	US 5579468-A 4-25-Nov-1996;						
FEATURES	/organism="unknown"						
BASE COUNT	3C a 1 C 0 g 0 t						
ORIGIN							
Query Match							
Best Local Similarity	83.3%						
Matches	37;						
DEFINITION	Linear						
ACCESION	129931						
VERSION	GI:1820722						
KEYWORDS							
SOURCE							
ORGANISM							
REFERENCE	1 (bases 1 to 37)						
AUTHORS							
TITLE							
JOURNAL							
FEATURES	/organism="unknown"						
BASE COUNT	2 a 18 C 0 g 18 t						
ORIGIN							
Query Match							
Best Local Similarity	81.6%						
Matches	31;						
DEFINITION	Conservative						
ACCESION	0; Mismatches						
VERSION	7; Indels						
KEYWORDS	0; Gaps						
SOURCE	0;						
ORGANISM							
REFERENCE	1 (bases 1 to 38)						
AUTHORS	Good, A. G.						
TITLE	Tissue-specific expression of target genes in plants						
JOURNAL	Patent - WO 9115633-A 6-02-AUG-2001;						
FEATURES	THE GOVERNORS OF THE UNIVERSITY OF ALBERTA (CA)						
BASE COUNT	1..38						
ORIGIN	/organism="Brassica napus"						
Query Match							
Best Local Similarity	81.6%						
Matches	31;						
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ACCESION	0; Mismatches						
VERSION	7; Indels						
KEYWORDS	0; Gaps						
SOURCE	0;						
ORGANISM							
REFERENCE	1 (bases 1 to 38)						
AUTHORS	Obara, M., Ohara, K., Tabira, K., Matsuzaki, T. and Ono, H.						
TITLE	Vector expressing full length gene of RNA virus and utilization						
JOURNAL	Patent - JP 2000-275303-A 19-06-JUN-2000;						
FEATURES	TRIZYU MEITA-ULTRA ORGANIZATION FIP MEDICAL RESEARCH, CHIGAI						
BASE COUNT	2 a 18 C 0 g 18 t						
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Query Match							
Best Local Similarity	81.6%						
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DEFINITION	Conservative						
ACCESION	0; Mismatches						
VERSION	7; Indels						
KEYWORDS	0; Gaps						
SOURCE	0;						
ORGANISM							
REFERENCE	1 (bases 1 to 38)						
AUTHORS	Obara, M., Ohara, K., Tabira, K., Matsuzaki, T. and Ono, H.						
TITLE	Vector expressing full length gene of RNA virus and utilization						
JOURNAL	Patent - JP 2000-275303-A 19-06-JUN-2000;						
FEATURES	TRIZYU MEITA-ULTRA ORGANIZATION FIP MEDICAL RESEARCH, CHIGAI						
BASE COUNT	2 a 18 C 0 g 18 t						
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Query Match							
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Matches	31;						
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ACCESION	0; Mismatches						
VERSION	7; Indels						
KEYWORDS	0; Gaps						
SOURCE	0;						
ORGANISM							
REFERENCE	1 (bases 1 to 38)						
AUTHORS	Obara, M., Ohara, K., Tabira, K., Matsuzaki, T. and Ono, H.						
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JOURNAL	Patent - JP 2000-275303-A 19-06-JUN-2000;						
FEATURES	TRIZYU MEITA-ULTRA ORGANIZATION FIP MEDICAL RESEARCH, CHIGAI						
BASE COUNT	2 a 18 C 0 g 18 t						
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Query Match							
Best Local Similarity	81.6%						
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DEFINITION	Conservative						
ACCESION	0; Mismatches						
VERSION	7; Indels						
KEYWORDS	0; Gaps						
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ORGANISM							
REFERENCE	1 (bases 1 to 38)						
AUTHORS	Obara, M., Ohara, K., Tabira, K., Matsuzaki, T. and Ono, H.						
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JOURNAL	Patent - JP 2000-275303-A 19-06-JUN-2000;						
FEATURES	TRIZYU MEITA-ULTRA ORGANIZATION FIP MEDICAL RESEARCH, CHIGAI						
BASE COUNT	2 a 18 C 0 g 18 t						
ORIGIN							
Query Match							
Best Local Similarity	81.6%						
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DEFINITION	Conservative						
ACCESION	0; Mismatches						
VERSION	7; Indels						
KEYWORDS	0; Gaps						
SOURCE	0;						
ORGANISM							
REFERENCE	1 (bases 1 to 38)						
AUTHORS	Obara, M., Ohara, K., Tabira, K., Matsuzaki, T. and Ono, H.						
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JOURNAL	Patent - JP 2000-275303-A 19-06-JUN-2000;						
FEATURES	TRIZYU MEITA-ULTRA ORGANIZATION FIP MEDICAL RESEARCH, CHIGAI						
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ACCESION	0; Mismatches						
VERSION	7; Indels						
KEYWORDS	0; Gaps						
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ORGANISM							
REFERENCE	1 (bases 1 to 38)						
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JOURNAL	Patent - JP 2000-275303-A 19-06-JUN-2000;						
FEATURES	TRIZYU MEITA-ULTRA ORGANIZATION FIP MEDICAL RESEARCH, CHIGAI						
BASE COUNT	2 a 18 C 0 g 18 t						
ORIGIN							
Query Match							
Best Local Similarity	81.6%						
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VERSION	7; Indels						
KEYWORDS	0; Gaps						
SOURCE	0;						
ORGANISM							
REFERENCE	1 (bases 1 to 38)						
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JOURNAL	Patent - JP 2000-275303-A 19-06-JUN-2000;						
FEATURES	TRIZYU MEITA-ULTRA ORGANIZATION FIP MEDICAL RESEARCH, CHIGAI						
BASE COUNT	2 a 18 C 0 g 18 t						
ORIGIN							
Query Match							
Best Local Similarity	81.6%						
Matches	31;						
DEFINITION	Conservative						
ACCESION	0; Mismatches						
VERSION	7; Indels						
KEYWORDS	0; Gaps						
SOURCE	0;						
ORGANISM							
REFERENCE	1 (bases 1 to 38)						
AUTHORS	Obara, M., Ohara, K., Tabira, K., Matsuzaki, T. and Ono, H.						
TITLE	Vector expressing full length gene of RNA virus and utilization						
JOURNAL	Patent - JP 2000-275303-A 19-06-JUN-2000;						
FEATURES	TRIZYU MEITA-ULTRA ORGANIZATION FIP MEDICAL RESEARCH, CHIGAI						
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ORIGIN							
Query Match							
Best Local Similarity	81.6%						
Matches	31;						
DEFINITION	Conservative						
ACCESION	0; Mismatches						
VERSION	7; Indels						
KEYWORDS	0; Gaps						
SOURCE	0;						
ORGANISM							
REFERENCE	1 (bases 1 to 38)						
AUTHORS	Obara, M., Ohara, K., Tabira, K., Matsuzaki, T. and Ono, H.						
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JOURNAL	Patent - JP 2000-275303-A 19-06-JUN-2000;						
FEATURES	TRIZYU MEITA-ULTRA ORGANIZATION FIP MEDICAL RESEARCH, CHIGAI						
BASE COUNT	2 a 18 C 0 g 18 t						
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Query Match							
Best Local Similarity	81.6%						
Matches	31;						
DEFINITION	Conservative						
ACCESION	0; Mismatches						
VERSION	7; Indels						
KEYWORDS	0; Gaps						
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ORGANISM							
REFERENCE	1 (bases 1 to 38)						
AUTHORS	Obara, M., Ohara, K., Tabira, K., Matsuzaki, T. and Ono, H.						
TITLE	Vector expressing full length gene of RNA virus and utilization						
JOURNAL	Patent - JP 2000-275303-A 19-06-JUN-2000;						
FEATURES	TRIZYU MEITA-ULTRA ORGANIZATION FIP MEDICAL RESEARCH, CHIGAI						
BASE COUNT	2 a 18 C 0 g 18 t						
ORIGIN							
Query Match			</td				





DEFINITION Sequence 5 from patent US 5571893 .

VERSION I28514  
KEYWORDS Unknown.

ORGANISM Unknown.

REFERENCE 1 (bases 1 to 50)  
AUTHORS Baker,J., Chien,K., King,K., Pennica,D. and Wood,W.  
TITLE JOURNAL Patent: US 5571893-A 5 NOV-1995;  
FEATURES Location/Qualifiers  
source 1..50  
/organism="unknown"  
BASE COUNT 3 a 7 c 7 g 33 t

ORIGIN Query Match Score 26.2%; DB 6; Length 50;  
Best Local Similarity 72.3%; Pred. No. 6.7e+05;  
Matches 34; Conservative 0; Mismatches 13; Indels 0; Gaps 0;

Qy 1240 AAAAAAACAAACAAAAACCCAAAGTCCCCCTCCGGCAG 1286  
Db 48 AAAAAAAAATTCGAGCTCCGGCG 2

RESULT 32  
I70295/c  
LOCUS 170295 Sequence 5 from patent US 5679545.  
DEFINITION 50 bp DNA  
ACCESSION 170295  
VERSION 170295 1 GI:30066430  
KEYWORDS Unknown.  
SOURCE Unknown.  
ORGANISM Unknown.  
REFERENCE 1 (bases 1 to 50)  
AUTHORS Baker,J., Chien,K., King,K., Pennica,D. and Wood,W.  
TITLE Gene encoding cardiac hypertrophy factor  
JOURNAL Patent: US 5679545-A 5 21-OCT-1997;  
FEATURES Location/Qualifiers  
source 1..50  
/organism="unknown"  
BASE COUNT 3 a 7 c 7 g 33 t

ORIGIN Query Match Score 26.2%; DB 6; Length 50;  
Best Local Similarity 72.3%; Pred. No. 6.7e+05;  
Matches 34; Conservative 0; Mismatches 13; Indels 0; Gaps 0;

Qy 1240 AAAAAAACAAACAAAAACCCAAAGTCCCCCTCCGGCAG 1286  
Db 48 AAAAAAAAATTCGAGCTCCGGCG 2

RESULT 30  
I41125/c  
LOCUS 141125 Sequence 5 from patent US 5624806.  
DEFINITION 50 bp DNA  
VERSION I41125  
KEYWORDS Unknown.  
ORGANISM Unknown.  
REFERENCE 1 (bases 1 to 50)  
AUTHORS Baker,J., Chien,K., King,K., Pennica,D. and Wood,W.  
TITLE Antibodies to cardiac hypertrophy factor and uses thereof  
JOURNAL Patent: US 5624806-A 5 29-APR-1997;  
FEATURES Location/Qualifiers  
source 1..50  
/organism="unknown"  
BASE COUNT 3 a 7 c 7 g 33 t

ORIGIN Query Match Score 26.2%; DB 6; Length 50;  
Best Local Similarity 72.3%; Pred. No. 6.7e+05;  
Matches 34; Conservative 0; Mismatches 13; Indels 0; Gaps 0;

Qy 1240 AAAAAAACAAACAAAAACCCAAAGTCCCCCTCCGGCAG 1286  
Db 48 AAAAAAAAATTCGAGCTCCGGCG 2

RESULT 31  
I49056/c  
LOCUS 149056 Sequence 5 from patent US 5627073.  
DEFINITION 50 bp DNA  
VERSION I49056  
KEYWORDS Unknown.  
ORGANISM Unknown.  
REFERENCE 1 (bases 1 to 50)  
AUTHORS Baker,J., Chien,K., King,K., Pennica,D. and Wood,W.  
TITLE Hybrids producing antibodies to cardiac hypertrophy factor  
JOURNAL Patent: US 5627073-A 5 06-MAY-1997;  
FEATURES Location/Qualifiers  
source 1..50  
/organism="unknown"  
BASE COUNT 3 a 7 c 7 g 33 t

ORIGIN Query Match Score 26.2%; DB 6; Length 50;  
Best Local Similarity 72.3%; Pred. No. 6.7e+05;  
Matches 34; Conservative 0; Mismatches 13; Indels 0; Gaps 0;

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Db 48 AAAAAAAAATTCGAGCTCCGGCG 2

RESULT 34  
AX458031/c  
LOCUS AX458031 Sequence 23 from Patent WO0246387.  
DEFINITION 47 bp DNA  
BASE COUNT 33 t

ORIGIN Query Match Score 26.2%; DB 6; Length 50;  
Best Local Similarity 72.3%; Pred. No. 6.7e+05;  
Matches 34; Conservative 0; Mismatches 13; Indels 0; Gaps 0;

Qy 1240 AAAAAAACAAACAAAAACCCAAAGTCCCCCTCCGGCAG 1286  
Db 48 AAAAAAAAATTCGAGCTCCGGCG 2





